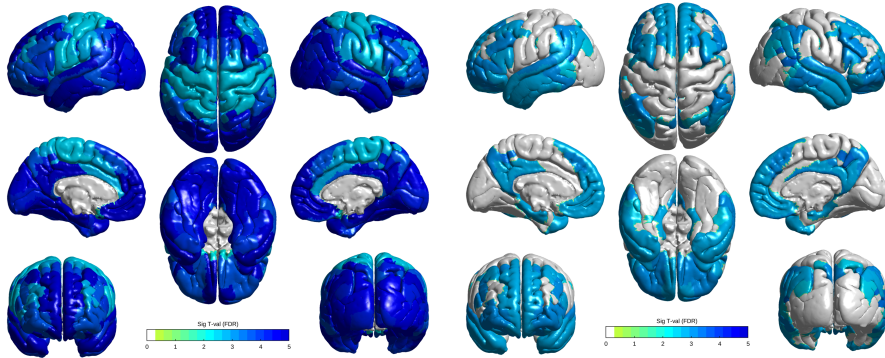
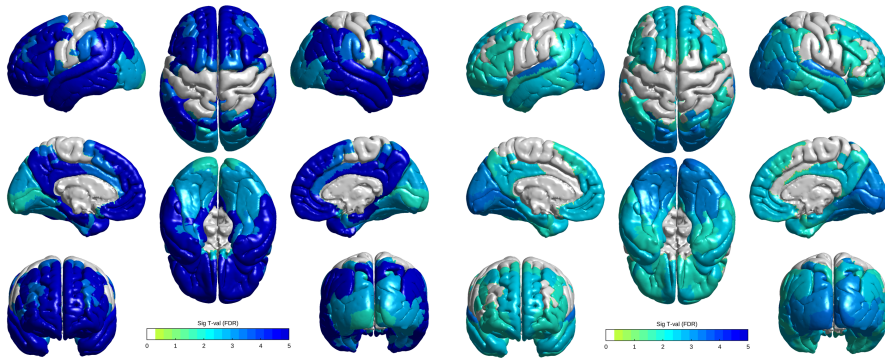


Supplemental Figure 1. Flowchart of subject inclusion in primary and secondary analyses. ADNI1, ADNI-GO and ADNI2 records for individuals with MRI, FDG-PET and baseline CSF biomarker analyses were included. A total of 887 subjects were used in primary analyses of baseline neuroimaging measures, while 816 were used in secondary analyses where a month 12 follow-up cognition visit was required.

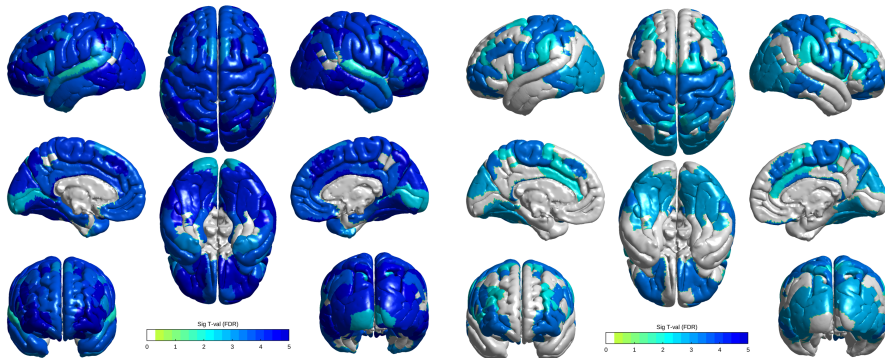
AD Continuum v. SNAP



A. FDG-PET Hypometabolism



B. Cortical Atrophy



C. Network Concordance

Supplemental Figure 2. T-Scores of FDR-corrected significant networks in AD-Continuum (N = 559) as compared to SNAP (N = 149) in W-score transformed Cortical Thickness (top panel), FDG-PET hypometabolism (middle panel) and Network Concordance (bottom panel) after controlling for sex, education, APOE4 (left panel), as well as CDR-SB (right panel).